

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: February 26, 2002, 11:47:52 : Search time 3596.29 Seconds
(without alignments)
3318.831 Million cell updates/sec

Title: US-09-602-833a-3
Perfect score: 681
Sequence: 1 atgagaattcgtatctgcgc.....cttttagccttcacttga 681

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapept 1.0

Searched: 17159718 seqs, 876320856 residues

Total number of hits satisfying chosen parameters: 34319436

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Pending_Patents_NA_Main:*

1: /cgn2_6/ptodata/2/pna/US06_COMB.seq:*
2: /cgn2_6/ptodata/2/pna/US06_COMB.seq:*
3: /cgn2_6/ptodata/2/pna/US07_COMB.seq:*
4: /cgn2_6/ptodata/2/pna/US08_COMB.seq:*
5: /cgn2_6/ptodata/2/pna/US08_COMB.seq:*
6: /cgn2_6/ptodata/2/pna/US08_COMB.seq:*
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10: /cgn2_6/ptodata/2/pna/US08_COMB.seq:*
11: /cgn2_6/ptodata/2/pna/US08_COMB.seq:*
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43: /cgn2_6/ptodata/2/pna/US09_COMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	681	100.0	681	47	US-60-140-627-3
2	681	100.0	1116	1	PCT-US01-14826-12
3	681	100.0	1989	1	PCT-US01-14826-12
4	681	100.0	2094	26	US-09-667-298-12
5	661.8	97.2	2035	1	PCT-US01-14826-1444
6	661.8	97.2	2035	22	US-09-577-408-1843
7	632	92.8	2010	23	US-09-616-111-806
8	632	92.8	2010	25	US-09-652-915-2927
9	576	84.6	577	19	US-60-245-228-444
10	504.8	74.1	630	19	US-09-516-448-1391
11	504.8	74.1	630	31	US-09-833-181-1391
12	372.2	54.7	484	23	US-09-616-111-636
13	345	50.7	840	26	US-09-668-683-3967
14	345	50.7	840	26	US-60-168-611-2255
15	324.4	47.6	327	17	US-09-332-782-6811
16	324.4	47.6	327	19	US-09-515-694-6811
17	302.6	44.4	384	12	US-09-915-738-10130
18	302.6	44.4	384	32	US-09-915-738-10130
19	265	38.9	334	17	US-09-359-067-5831
20	195.6	28.7	323	12	US-08-824-056-1404
21	187.4	27.5	239	14	US-09-075-782-603
22	187.4	27.5	239	21	US-09-540-766-17283
23	187.4	27.5	239	37	US-60-046-624-603
24	172	25.3	480	32	US-09-933-524-29336
25	172	25.3	480	32	US-09-933-524-29336
26	161	23.6	15011	57	US-60-245-228-24
27	128	18.8	440	19	US-09-528-409-25490
28	128	18.8	440	32	US-09-933-524-25490
29	125	18.4	456	19	US-09-528-409-24337
30	125	18.4	456	32	US-09-933-524-24337
31	118.4	17.4	494	52	US-60-195-050-71
32	117.2	17.2	222	23	US-09-619-303-42
33	107.8	15.8	1660	25	US-09-649-164-9252
34	107.8	15.8	1660	29	US-09-726-802-2039
35	107.8	15.8	1660	29	US-09-726-802-2039
36	107.8	15.8	1856	25	US-09-652-913-8543
37	107.8	15.8	2122	31	US-09-516-448-1392
38	107.8	15.8	2122	31	US-09-833-381-1392
39	100	14.7	1434	18	US-09-496-914A-2777
40	100	14.7	1434	22	US-09-560-875A-2777
41	98.4	14.4	1848	17	US-09-359-922-5894
42					
43					

42	98.4	14.4	1848	17	US-09-359-922-5894	Sequence 5894, Ap
43	87	12.8	429	19	US-09-528-409-13035	Sequence 13035, A
44	87	12.8	429	32	US-09-933-524-13035	Sequence 13035, A
45	85	12.5	823	57	US-60-245-228-523	Sequence 523, App

ALIGNMENTS

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RESULT 1
US-60-140-627-3
; Sequence 3, Application US/60140627
; GENERAL INFORMATION:
; APPLICANT: Turner, Alex
; APPLICANT: Zambrowicz, Brian
; APPLICANT: Nehls, Michael
; APPLICANT: Freidrich, Glenn A.
; APPLICANT: Sands, Arthur T.
; TITLE OF INVENTION: A NOVEL HUMAN CDNA CLONE AND PROTEINS
; TITLE OF INVENTION: ENCODED THEREBY
; FILE REFERENCE: 8535-0036-888
; CURRENT APPLICATION NUMBER: US/60/140, 627
; CURRENT FILING DATE: 1999-06-23
; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 3
; LENGTH: 681
; TYPE: DNA
; ORGANISM: Homo sapien
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)...(678)
US-60-140-627-3
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Query Match Best Local Similarity 100.0%; Score 681; DB 47; Length 681;
Matches 681; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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QY 1 atgagaattcgtatctgcacaaacccaatctcacatcttcacagcagaatcggtgt 60
Db 1 atgagaattcgtatctgcacaaacccaatctcacatcttcacagcagaatcggtgt 60
QY 61 ttgaagaacctgaaagaaactcaatgtggttcaactatctgaagaagcattccctcagaa 120
Db 61 ttgaagaacctgaaagaaactcaatgtggttcaactatctgaagaagcattccctcagaa 120
QY 121 ttggagagctgtgaaactctagagagactgtgtcttggaatctagaattatagag 180
Db 121 ttggagagctgtgaaactctagagagactgtgtcttggaatctagaattatagag 180
QY 181 ctgaccttgataagtaatttgaaagcaagttacatttgatagatctcagcaaacag 240
Db 181 ctgaccttgataagtaatttgaaagcaagttacatttgatagatctcagcaaacag 240
QY 241 ttcccaagtgtcccaatctgttcctcggaatgcgaattgcagtggttgatatacagc 300
Db 241 ttcccaagtgtcccaatctgttcctcggaatgcgaattgcagtggttgatatacagc 300
QY 301 agcaataacccgacgcagctgcgcaagatatagacaagctagagagctgcagaagctt 360
Db 301 agcaataacccgacgcagctgcgcaagatatagacaagctagagagctgcagaagctt 360
QY 361 ctctgtataaaaacaagttaacctacattccctatccatctgcagtgaactgaagaagctc 420
Db 361 ctctgtataaaaacaagttaacctacattccctatccatctgcagtgaactgaagaagctc 420
QY 421 actctgtatgcgtcagtgaggacatttggttgagagctcccaactgaccttgtaactca 480
Db 421 actctgtatgcgtcagtgaggacatttggttgagagctcccaactgaccttgtaactca 480
QY 481 tccacaccttaaatattgtaagccttatgacacatcttatgtaagcccaatgtgaa 540
Db 481 tccacaccttaaatattgtaagccttatgacacatcttatgtaagcccaatgtgaa 540
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Db 481 tccacaccttaaatattgtaagccttatgacacatcttatgtaagcccaatgtgaa 540
QY 541 gatggcaatgaaataatgaaagtgaaacggaatcgcacacatttgataaagaattag 600
Db 541 gatggcaatgaaataatgaaagtgaaacggaatcgcacacatttgataaagaattag 600
QY 601 aaagcctatatgaaagaccttaagaagaagaatctgttccagctatacccaagtg 660
Db 601 aaagcctatatgaaagaccttaagaagaagaatctgttccagctatacccaagtg 660
QY 661 tctttaagcctcaacttga 681
Db 661 tctttaagcctcaacttga 681
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RESULT 2
US-60-140-627-1
; Sequence 1, Application US/60140627
; GENERAL INFORMATION:
; APPLICANT: Turner, Alex
; APPLICANT: Zambrowicz, Brian
; APPLICANT: Nehls, Michael
; APPLICANT: Freidrich, Glenn A.
; APPLICANT: Sands, Arthur T.
; TITLE OF INVENTION: A NOVEL HUMAN CDNA CLONE AND PROTEINS
; TITLE OF INVENTION: ENCODED THEREBY
; FILE REFERENCE: 8535-0036-888
; CURRENT APPLICATION NUMBER: US/60/140, 627
; CURRENT FILING DATE: 1999-06-23
; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 1
; LENGTH: 1116
; TYPE: DNA
; ORGANISM: Homo sapien
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)...(1113)
US-60-140-627-1
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Query Match Best Local Similarity 100.0%; Score 681; DB 47; Length 1116;
Matches 681; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 1 atgagaattcgtatctgcacaaacccaatctcacatcttcacagcagaatcggtgt 60
QY 436 atgagaattcgtatctgcacaaacccaatctcacatcttcacagcagaatcggtgt 495
Db 436 atgagaattcgtatctgcacaaacccaatctcacatcttcacagcagaatcggtgt 495
QY 61 ttgaagaacctgaaagaaactcaatgtggttcaactatctgaagaagcattccctcagaa 120
Db 61 ttgaagaacctgaaagaaactcaatgtggttcaactatctgaagaagcattccctcagaa 120
QY 121 ttggagagctgtgaaactctagagagactgtgtcttggaatctagaattatagag 180
Db 121 ttggagagctgtgaaactctagagagactgtgtcttggaatctagaattatagag 180
QY 181 ctgaccttgataagtaatttgaaagcaagttacatttgatagatctcagcaaacag 240
Db 181 ctgaccttgataagtaatttgaaagcaagttacatttgatagatctcagcaaacag 240
QY 241 ttcccaagtgtcccaatctgttcctcggaatgcgaattgcagtggttgatatacagc 300
Db 241 ttcccaagtgtcccaatctgttcctcggaatgcgaattgcagtggttgatatacagc 300
QY 301 agcaataacccgacgcagctgcgcaagatatagacaagctagagagctgcagaagctt 360
Db 301 agcaataacccgacgcagctgcgcaagatatagacaagctagagagctgcagaagctt 360
QY 361 ctctgtataaaaacaagttaacctacattccctatccatctgcagtgaactgaagaagctc 420
Db 361 ctctgtataaaaacaagttaacctacattccctatccatctgcagtgaactgaagaagctc 420
QY 796 ctctgtataaaaacaagttaacctacattccctatccatctgcagtgaactgaagaagctc 855
Db 796 ctctgtataaaaacaagttaacctacattccctatccatctgcagtgaactgaagaagctc 855
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QY 421 actctgtatgctgcgaagtgaggaccatttggtgagctcccaactgaccttctgtactca 480
Db 826 actctgtatgctgcgaagtgaggaccatttggtgagctcccaactgaccttctgtactca 915
QY 481 tccacaccttaaaattgttaagccttatgacaacacctatgtataatgacccaatgtgaa 540
Db 916 tccacaccttaaaattgttaagccttatgacaacacctatgtataatgacccaatgtgaa 975
QY 541 gatggcaatgaataatgaagaatgaaagtgatgcgcgaaccttttggataaagaagtatg 600
Db 976 gatggcaatgaataatgaagaatgaaagtgatgcgcgaaccttttggataaagaagtatg 1035
QY 601 aaagcctatatgaagaccttaagaagaagaatctgttccgaagctataccaccaagaatg 660
Db 1036 aaagcctatatgaagaccttaagaagaagaatctgttccgaagctataccaccaagaatg 1095
QY 661 tcttttagccttcaactttga 681
Db 1096 tcttttagccttcaactttga 1116

RESULT 3
PCT-US01-14826-12
Sequence 12, Application PC/TUS0114826
GENERAL INFORMATION:
APPLICANT: Hyseq, Inc
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
FILE REFERENCE: 21272-103
CURRENT APPLICATION NUMBER: PCT/US01/14826
CURRENT FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: 09/577,408
PRIOR FILING DATE: 2000-05-18
PRIOR APPLICATION NUMBER: 09/677,298<151> 2000-09-22
PRIOR APPLICATION NUMBER: 09/695,781<151> 2000-10-24
PRIOR APPLICATION NUMBER: 09/715,869<151> 2000-11-17
PRIOR APPLICATION NUMBER: 09/775,330<151> 2001-02-01
NUMBER OF SEQ ID NOS: 864
SOFTWARE: Custom
SEQ ID NO 12
LENGTH: 1989
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (330)..(1442)
PCT-US01-14826-12

Query Match 100.0%; Score 681; DB 1; Length 1989;
Best Local Similarity 100.0%; Pred. NO. 6.3e-181;
Matches 681; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 atgagaattcgtgattctgcgaataaaacaaatctcacattctccagcagaatcggtgt 60
Db 765 atgagaattcgtgattctgcgaataaaacaaatctcacattctccagcagaatcggtgt 824
QY 61 ttgaagaacctgaagaacctcaatggtgttccaactatctgaagagacattccctccagaa 120
Db 825 ttgaagaacctgaagaacctcaatggtgttccaactatctgaagagacattccctccagaa 884
QY 121 ttgggagattgtgaaaatctagagagactgattgttctggaatctagaattaaatgag 180
Db 885 ttgggagattgtgaaaatctagagagactgattgttctggaatctagaattaaatgag 944
QY 181 ctgcaccttgtaataatgttaagaagttacatttgaatatactccgcaacaag 240
Db 945 ctgcaccttgtaataatgttaagaagttacatttgaatatactccgcaacaag 1004
QY 241 ttcccaatgtcccaaatgtgtccctgcgagatgcgaatttgcgaatgtgttgatatcagc 300
Db 1005 ttcccaatgtcccaaatgtgtccctgcgagatgcgaatttgcgaatgtgttgatatcagc 1064
QY 301 agcaataacctgaacctctgcgcaagataatagacaggtagaagctgtcagagctt 360

Db 1065 agcaataacctgaacctctgcgcaagataatagacaggtagaagctgtcagagctt 1124
QY 361 ctctgtataaaacaagtgtacctacccttccattccatgctgcaacctgaagaagctc 420
Db 1125 ctctgtataaaacaagtgtacctacccttccattccatgctgcaacctgaagaagctc 1184
QY 421 actctgtatgctgcgaagtgaggaccatttggtgagctcccaactgaccttctgtactca 480
Db 1185 actctgtatgctgcgaagtgaggaccatttggtgagctcccaactgaccttctgtactca 1244
QY 481 tccacaccttaaaattgttaagccttatgacaacacctatgtataatgacccaatgtgaa 540
Db 1245 tccacaccttaaaattgttaagccttatgacaacacctatgtataatgacccaatgtgaa 1304
QY 541 gatggcaatgaataatgaagaatgaaagtgatgcgcgaaccttttggataaagaagtatg 600
Db 1305 gatggcaatgaataatgaagaatgaaagtgatgcgcgaaccttttggataaagaagtatg 1364
QY 601 aaagcctatatgaagaccttaagaagaagaatctgttccgaagctataccaccaagaatg 660
Db 1365 aaagcctatatgaagaccttaagaagaagaatctgttccgaagctataccaccaagaatg 1424
QY 661 tcttttagccttcaactttga 681
Db 1425 tcttttagccttcaactttga 1445

RESULT 4
US-09-667-298-12
Sequence 12, Application US/09667298
GENERAL INFORMATION:
APPLICANT: Tang, Y. Tom
APPLICANT: Liu, Chenghua
APPLICANT: Zhou, Ping
APPLICANT: Asundi, Vinod
APPLICANT: Ren, Feiyan
APPLICANT: Zhao, Qing A.
APPLICANT: Zhang, Jie
APPLICANT: Xue, Aidong J.
APPLICANT: Wang, Jian-Rui
APPLICANT: Chen, Rui-hong
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: Novel Nucleic Acids and
FILE REFERENCE: 792CIP2A
CURRENT APPLICATION NUMBER: US/09/667,298
CURRENT FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: 09/577,408
PRIOR FILING DATE: 2000-05-18
NUMBER OF SEQ ID NOS: 178
SOFTWARE: pt_FL-genes Version 2.0
SEQ ID NO 12
LENGTH: 2094
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (435)..(1550)
US-09-667-298-12

Query Match 100.0%; Score 681; DB 26; Length 2094;
Best Local Similarity 100.0%; Pred. NO. 6.4e-181;
Matches 681; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 atgagaattcgtgattctgcgaataaaacaaatctcacattctccagcagaatcggtgt 60
Db 870 atgagaattcgtgattctgcgaataaaacaaatctcacattctccagcagaatcggtgt 929
QY 61 ttgaagaacctgaagaacctcaatggtgttccaactatctgaagagacattccctccagaa 120
Db 930 ttgaagaacctgaagaacctcaatggtgttccaactatctgaagagacattccctccagaa 989

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QY 121 ttggagagctgtgaaatctagagagactggtgttcttgynaactctagaattatgtgag 180
|||
Db 990 ttggagagctgtgaaatctagagagactggtgttcttgynaactctagaattatgtgag 1049
QY 181 ctgccccttgaaatgaatattgaaagcaagtatacttcttgatatacttcagcaacaag 240
|||
Db 1050 ctgccccttgaaatgaatattgaaagcaagtatacttcttgatatacttcagcaacaag 1109
QY 241 ttcttcagtggtcccaatctgtgtctctgagatggtcgaatttgcaatgtgtgttgatatacgc 300
|||
Db 1110 ttcttcagtggtcccaatctgtgtctctgagatggtcgaatttgcaatgtgtgttgatatacgc 1169
QY 301 agcaataaccctgaccgacctgctgcgaagatatagacagctagagagctgagagctt 360
|||
Db 1170 agcaataaccctgaccgacctgctgcgaagatatagacagctagagagctgagagctt 1229
QY 361 ctcttgataaaaaaagaattgacacttccctatccatgctgtaacctgaaagagctc 420
|||
Db 1230 ctcttgataaaaaaagaattgacacttccctatccatgctgtaacctgaaagagctc 1289
QY 421 actctgttagtgcgtcagtgaggagaccattgtgtgagctcccaactgaccttgtgactca 480
|||
Db 1290 actctgttagtgcgtcagtgaggagaccattgtgtgagctcccaactgaccttgtgactca 1349
QY 481 tccacacctttaaattgtgtaagccttatggaacaatccatgtataatgcccgaatgtgaa 540
|||
Db 1350 tccacacctttaaattgtgtaagccttatggaacaatccatgtataatgcccgaatgtgaa 1409
QY 541 gatgcaatgnaatataatgaaagtgaaagtgacggaatgcgaacatttggataaagaattatg 600
|||
Db 1410 gatgcaatgnaatataatgaaagtgaaagtgacggaatgcgaacatttggataaagaattatg 1469
QY 601 aaagcctatattgaagaccttaaaagaagaatctgttccagctataccaccagaatg 660
|||
Db 1470 aaagcctatattgaagaccttaaaagaagaatctgttccagctataccaccagaatg 1529
QY 661 tcttttagcctcaacttga 681
|||
Db 1530 tcttttagcctcaacttga 1550

RESULT 5
PCT-US01-14826-444
; Sequence 444, Application PC/TUS0114826
; GENERAL INFORMATION:
; APPLICANT: Hyseq, Inc
; TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
; FILE REFERENCE: 21272-103
; CURRENT APPLICATION NUMBER: PCT/US01/14826
; CURRENT FILING DATE: 2001-05-16
; PRIOR APPLICATION NUMBER: 09/577,408
; PRIOR FILING DATE: 2000-05-18
; PRIOR APPLICATION NUMBER: 09/677,298<151> 2000-09-22
; PRIOR APPLICATION NUMBER: 09/695,781<151> 2000-10-24
; PRIOR APPLICATION NUMBER: 09/715,869<151> 2000-11-17
; PRIOR APPLICATION NUMBER: 09/775,330<151> 2001-02-01
; NUMBER OF SEQ ID NOS: 864
; SOFTWARE: Custom
; SEQ ID NO 444
; LENGTH: 2035
; TYPE: DNA
; ORGANISM: Homo sapiens
PCT-US01-14826-444
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Query Match 97.2%; Score 661.8; DB 1; Length 2035;
Best Local Similarity 96.2%; Pred. NO. 1.6e-175;
Matches 669; Conservative 0; Mismatches 12; Indels 0; Gaps 0;
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QY 1 atgagaattcgtgattccaaaacaaatctcacatcttcacagagaatcggtgt 60
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Db 811 atgagaattcgtgattccaaaacaaatctcacatcttcacagagaatcggtgt 870
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QY 61 ttgaagaacctgaaagaaatccaatgtggtttcacaatctgaagagattccctccagaa 120
|||
Db 871 ttgaagaacctgaaagaaatccaatgtggtttcacaatctgaagagattccctccagaa 930
QY 121 ttggagagattgtgaaatcttagagagactggtatttcttgaaactctagaattatgtgag 180
|||
Db 931 ttggagagattgtgaaatcttagagagactggtatttcttgaaactctagaattatgtgag 990
QY 181 ctgccccttgaaatgaatattgaaagcaagtatacttcttgatatacttcagcaacaag 240
|||
Db 991 ctgccccttgaaatgaatattgaaagcaagtatacttcttgatatacttcagcaacaag 1050
QY 241 ttcttcagtggtcccaatctgtgtctctgagatggtcgaatttgcaatgtgtgttgatatacgc 300
|||
Db 1051 ttcttcagtggtcccaatctgtgtctctgagatggtcgaatttgcaatgtgtgttgatatacgc 1110
QY 301 agcaataaccctgaccgacctgctgcgaagatatagacagctagagagctgagagctt 360
|||
Db 1111 agcaataaccctgaccgacctgctgcgaagatatagacagctagagagctgagagctt 1170
QY 361 ctcttgataaaaaaagaattgacacttccctatccatgctgtaacctgaaagagctc 420
|||
Db 1171 ctcttgataaaaaaagaattgacacttccctatccatgctgtaacctgaaagagctc 1230
QY 421 actctgttagtgcgtcagtgaggagaccattgtgtgagctcccaactgaccttgtgactca 480
|||
Db 1231 actctgttagtgcgtcagtgaggagaccattgtgtgagctcccaactgaccttgtgactca 1290
QY 481 tccacacctttaaattgtgtaagccttatggaacaatccatgtataatgcccgaatgtgaa 540
|||
Db 1291 tccacacctttaaattgtgtaagccttatggaacaatccatgtataatgcccgaatgtgaa 1350
QY 541 gatgcaatgnaatataatgaaagtgaaagtgacggaatgcgaacatttggataaagaattatg 600
|||
Db 1351 gatgcaatgnaatataatgaaagtgaaagtgacggaatgcgaacatttggataaagaattatg 1410
QY 601 aaagcctatattgaagaccttaaaagaagaatctgttccagctataccaccagaatg 660
|||
Db 1411 aaagcctatattgaagaccttaaaagaagaatctgttccagctataccaccagaatg 1470
QY 661 tcttttagcctcaacttga 681
|||
Db 1471 tcttttagcctcaacttga 1491
```

```
RESULT 6
US-09-577-408-1843
; Sequence 1843, Application US/09577408
; GENERAL INFORMATION:
; APPLICANT: Tang, Y. Tom
; APPLICANT: Tillinghast, John
; APPLICANT: Sinku, Ankura
; APPLICANT: Liu, Chenghua
; APPLICANT: Drmanac, Radoje T.
; TITLE OF INVENTION: Novel Nucleic Acids and
; FILE REFERENCE: 792
; CURRENT APPLICATION NUMBER: US/09/577,408
; CURRENT FILING DATE: 2000-05-18
; NUMBER OF SEQ ID NOS: 8502
; SOFTWARE: PL_CCT_genes Version 1.0
; SEQ ID NO 1843
; LENGTH: 2035
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (562)...(1446)
; OTHER INFORMATION: similar to g14538934 in the genepept database release 115,
; OTHER INFORMATION: Run with FASTX 3.3000, default parameters
US-09-577-408-1843
```

```

Query Match 97.2% Score 661.8; Db 22; Length 2035;
Best Local Similarity 98.2%; Prd. No. 1, 6e-175;
Matches 669; Conservative 0; Mismatches 12; Indels 0; Gaps 0;

QY 1 atggaattctgatactctgccaanaaacccaatctcaatcttccagcagaatctgtgt 60
Db 811 atggaattctgatactctgccaanaaacccaatctcaatcttccagcagaatctgtgt 870
QY 61 ttgaagaacctgaaagaactaactatgtgtttcaactatcttgaagaacttctccgaa 120
Db 871 ttgaagaacctgaaagaactaactatgtgtttcaactatcttgaagaacttctccgaa 930
QY 121 ttggagatctgaaatctagagagactgattgttctggaatctcagaattaatgag 180
Db 931 ttggagatctgaaatctagagagactgattgttctggaatctcagaattaatgag 990
QY 181 ctgcctcttgaattaagtaatttgaagcaagttacatttgtagatcttcagcaaacag 240
Db 991 ctgcctcttgaattaagtaatttgaagcaagttacatttgtagatcttcagcaaacag 1050
QY 241 ttccacagtcgccaactctgtctctcgagatgfcggaattgcagtggttgatcagc 300
Db 1051 ttccacagtcgccaactctgtctctcgagatgfcggaattgcagtggttgatcagc 1110
QY 301 agcaataactgacacgcagcttcgcgaagatagacagcgttagagagctgcagagctt 360
Db 1111 agcaataactgacacgcagcttcgcgaagatagacagcgttagagagcgttcagagctt 1170
QY 361 ctcttgtataaaaacaagttgaactaccttccactatccatgctgaacctgaagaagctc 420
Db 1171 ctcttgtataaaaacaagttgaactaccttccactatccatgctgaacctgaagaagctc 1230
QY 421 acctgtgtaagtcgacgaatgtaggacatttgttgaagcttccaaatgccttcttgatca 480
Db 1231 acctgtgtaagtcgacgaatgtaggacatttgttgaagcttccaaatgccttcttgatca 1290
QY 481 tccacacctttaaatttgttaagccttaaggacatcttaattgaataatgccaagttaa 540
Db 1291 tccacacctttaaatttgttaagccttaaggacatcttaattgaataatgccaagttaa 1350
QY 541 gatggcaatgaaataatggaagaagtgaacggagatgcgcacaacatttgttaagaagaatc 600
Db 1351 gatggcaatgaaataatggaagaagtgaacggagatgcgcacaacatttgttaagaagaatc 1410
QY 601 aaagccatatttgaagaccttaagaagaagaatctgttccagacatataccacaagt 660
Db 1411 aaagccatatttgaagaccttaagaagaagaatctgttccagacatataccacaagt 1470
QY 661 tctttagccttcaacttga 681
Db 1471 tctttagccttcaacttga 1491

RESULT 7
US-09-616-111-806
: Sequence 806, Application US/09616111
: GENERAL INFORMATION:
: APPLICANT: Holtzman, Douglas A.
: TITLE OF INVENTION: NOCULEIC ACID MOLECULES DERIVED FROM A
: FILE OF INVENTION: HUMAN PITUITARY LIBRARY
: FILE REFERENCE: 1600.1142-001
: CURRENT APPLICATION NUMBER: US/09/616, 111
: CURRENT FILING DATE: 2000-07-13
: PRIOR APPLICATION NUMBER: 60/143, 618
: PRIOR FILING DATE: 1999-07-13
: NUMBER OF SEQ ID NOS: 816
: SOFTWARE: FastSeq for Windows Version 3.0
: SEQ ID NO 806
: LENGTH: 2010
: TYPE: DNA
: ORGANISM: Homo sapiens

```

```

US-09-616-111-806

Query Match          92.8%, Score 632, DB 23, Length 2010;
Best Local Similarity 97.7%, Pred. NO. 4.2e-167;
Matches 666; Conservative 0; Mismatches 5; Indels 11; Gaps 2;

QY 1 atggaattctggatctgcacaaacccaacatctacatctctccagcagaatctggtt 60
Db 797 atggaattctggatctgcacaaacccaacatctacatctctccagcagaatctggtt 856
QY 61 ttgaagaacctggaagaacatcaatggtgttctcaactctgaagaagatctctcagaa 120
Db 857 ttgaagaacctggaagaacatcaatggtgttctcaactctgaagaagatctctcagaa 916
QY 121 ttggagatcttgaanaactagagagactggtattcttgaanaactgaattaatgag 180
Db 917 ttggagatcttgaanaactagagagactggtattcttgaanaactgaattaatgag 976
QY 181 ctgacctttgaatgaatgaatttgaagaagaattacatttgtagatatccagaacaag 240
Db 977 ctgacctttgaatgaagaat-----aaacatttgaatatccagaacaag 1026
QY 241 ttctccagtgcgcccaatctgtctcctgcgcgagatctgaatttgcagtgttggatalcagc 300
Db 1027 ttctccagtgcgcccaatctgtctcctgcgcgagatctgaatttgcagtgttggatalcagc 1086
QY 301 agcaataacctgacccgacccgcgcgaagatatagacagctagagagagctgcagagcttt 360
Db 1087 agcaataacctgacccgacccgcgcgaagatatagacagctagagagagctgcagagcttt 1146
QY 361 ctcttggtataaacaagaattgaacctactctccatttccatgctgaactgaagaagctc 420
Db 1147 ctcttggtataaacaagaattgaacctactctccatttccatgctgaactgaagaagctc 1206
QY 421 actctgttaagtcgtcaagtgaggagaccatttgtgtgagctccccaacgacctgtgtactca 480
Db 1207 actctgttaagtcgtcaagtgaggagaccatttgtgtgagctccccaacgacctgtgtactca 1266
QY 481 tccaca-ccttaaaatttgaagccttaagacacatccatttgaataatgcccgaatgtga 539
Db 1267 tccacaacctttaaatttgaagccttaagacacatccatttgaataatgcccgaatgtga 1326
QY 540 agatgaggcaatgaataatggaagaatggaacggatctgcacaacttttgtataagaagttat 599
Db 1327 agatgaggcaatgaataatggaagaatggaacggatctgcacaacttttgtataagaagttat 1386
QY 600 gaagcccatatttgaagccttaagaagaagaacatcttccagctataccacaagaat 659
Db 1387 gaagcccatatttgaagccttaagaagaagaacatcttccagctataccacaagaat 1446
QY 660 gtctttagccttcaacttga 681
Db 1447 gtctttagccttcaacttga 1468

RESULT      8
US-09-652-915-9227
; Sequence 9227, Application US/09652915
; GENERAL INFORMATION:
; APPLICANT: Faib, Dean R.
; TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
; TITLE OF INVENTION: THEREFOR
; FILE REFERENCE: 1600.1175-001
; CURRENT APPLICATION NUMBER: US/09/652,915
; CURRENT FILING DATE: 2000-08-31
; PRIOR APPLICATION NUMBER: 60/152,110
; PRIOR FILING DATE: 1999-08-31
; NUMBER OF SEQ ID NOS: 10311
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 9227
; LENGTH: 2010

```

TYPE: DNA
ORGANISM: Homo sapiens
US-09-652-915-9227

Query Match 92.8%; Score 632; DB 25; Length 2010;
Best Local Similarity 97.7%; Pred. No. 4.2e-167;
Matches 666; Conservative 0; Mismatches 5; Indels 11; Gaps 2;

```
QY 1 atgaagattctgagatctcccaaaaacccaatctcacatctctccagcagaatcgggtgt 60
    |||||
Db 797 atgaagattctgagatctcccaaaaacccaatctcacatctctccagcagaatcgggtgt 856
    |||||

QY 61 ttgaagacctgaaagaaactcaatggtggttcaactatctgaagagcattccctccagaa 120
    |||||
Db 857 ttgaagacctgaaagaaactcaatggtggttcaactatctgaagagcattccctccagaa 916
    |||||

QY 121 ttggaagattgtaaaatctgagagagactgattgttctggaactctgaatataatgag 180
    |||||
Db 917 ttggaagattgtaaaatctgagagagactgattgttctggaactctgaatataatgag 976
    |||||

QY 181 ctgaccttgaataagtaatttgaagcaagtlacattgttagatatctcaacaacaag 240
    |||||
Db 977 ctgaccttgaataagtaatttgaagcaagtlacattgttagatatctcaacaacaag 1026
    |||||

QY 241 ttctccagtgcccaatctgtgtcctgcgagatgtcgaatttgagtggttgatatcagc 300
    |||||
Db 1027 ttctccagtgcccaatctgtgtcctgcgagatgtcgaatttgagtggttgatatcagc 1086
    |||||

QY 301 agcaataaccgagaccgacctgcgcgaagatatagacagctgagagagctgagagcttt 360
    |||||
Db 1087 agcaataaccgagaccgacctgcgcgaagatatagacagctgagagagctgagagcttt 1146
    |||||

QY 361 ctctgtataaaaaaagctgacactaccttccctattccatgtctgaacctgaagaagctc 420
    |||||
Db 1147 ctctgtataaaaaaagctgacactaccttccctattccatgtctgaacctgaagaagctc 1206
    |||||

QY 421 actcgttagtcgtcagtgaggagaccattgtgtgagctcccaacgaccttgtgtacc 480
    |||||
Db 1207 actcgttagtcgtcagtgaggagaccattgtgtgagctcccaacgaccttgtgtacc 1266
    |||||

QY 481 tccaca-gccttaaaatttgaagcctatgacacatccatttgaataagccaatgtga 539
    |||||
Db 1267 tccaca-gccttaaaatttgaagcctatgacacatccatttgaataagccaatgtga 1326
    |||||

QY 540 agatgycgaatgaataatgaaagtgaacgagatcgccaacatttgaataaagaagtcat 599
    |||||
Db 1327 agatgycgaatgaataatgaaagtgaacgagatcgccaacatttgaataaagaagtcat 1386
    |||||

QY 600 gaaagccatattgaaagccttaagaagaagaatctgttccagctataccacaagt 659
    |||||
Db 1387 gaaagccatattgaaagccttaagaagaagaatctgttccagctataccacaagt 1446
    |||||

QY 660 gtccttagcctcaacttga 681
    |||||
Db 1447 gtccttagcctcaacttga 1468
    |||||
```

RESULT 9
US-60-245-228-444/C
Sequence 444, Application US/60245228
GENERAL INFORMATION:
APPLICANT: Beasley, Ellen
TITLE OF INVENTION: ISOLATED HUMAN CYCLASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN CYCLASE PROTEINS, AND USES
FILE REFERENCE: C1000878
CURRENT APPLICATION NUMBER: US/60/245, 228
CURRENT FILING DATE: 2000-11-03
NUMBER OF SEQ ID NOS: 630
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 444
LENGTH: 577

TYPE: DNA
ORGANISM: HUMAN
US-60-245-228-444

Query Match 84.6%; Score 576; DB 57; Length 577;
Best Local Similarity 100.0%; Pred. No. 1.6e-151;
Matches 576; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
QY 56 gttgttgaagaaacctgaagaacccaatgtggtgttcaactatctgaagagcattcc 115
    |||||
Db 577 gttgttgaagaaacctgaagaacccaatgtggtgttcaactatctgaagagcattcc 518
    |||||

QY 116 cagaattgagagattgtgaaatctagaagagactgattgttctggaactcagaat 175
    |||||
Db 517 CAGAAATGGGAGATGTGAAATCTAGAGAGACTGATTTGTCGAAATCTAGAAATTA 458
    |||||

QY 176 tggagctgaccttgaataagtaatttgaagcaagttacacttgttaattctcagaa 235
    |||||
Db 457 TGGAGCTGCCCTTGAATTAAGTAATTTGAAGCAAGTTACATTTGTAGATATCTCAGCA 398
    |||||

QY 236 acaagtttccagtgcccaatctgtcctgcgagatgtcgaatttgagtggttgata 295
    |||||
Db 397 ACAAGTTTCCAGTGTCCCAATCTGTGTCTGTGCGATTCGAAATTTGCAAGTGTGATA 338
    |||||

QY 296 tcaagcaataaacttgacagacctgcgcgaagatatagacagctagaagagctgaga 355
    |||||
Db 337 TCAGCAGCAATTAACCTGACCGCCGCAAGATATAGACAGGCTTAGAGGAGCTGCAGA 278
    |||||

QY 356 gcttctcttctataaaaaaagtgagacttaccctccattatccatgtctgaacctga 415
    |||||
Db 277 GCTTCTCTTGTATATAAACAAGTTGACCTTCCCTATATTCAGTGTGAACCTGAGAA 218
    |||||

QY 416 agctcactctgttagctgagtgaggacatttggtgagctcccaactgaccttgtg 475
    |||||
Db 217 AGCTCACTGTGTAGTGTGAGTGAGGACAAATTTGTGAGAGCTCCCACTGCCCTTGTG 158
    |||||

QY 476 actatccacaccttaaaatttgaagccttatgaaacatccatttgaataagccaat 535
    |||||
Db 157 ACTCATCCACACACCTTTAAATTTGTATGACCTTATGACAAATCTTATATATGCCCAT 98
    |||||

QY 536 gtaagagtgagcaatgaataatgaaagtgaacgagatcgccaacatttgaataaaga 595
    |||||
Db 97 GTGAAGATGGCAATGAATATGAAAGTGAAGCGGATCGCCACATTTGTGATAAGAG 38
    |||||

QY 596 ttatgaagcctatatgaaagccttaagaagaag 631
    |||||
Db 37 TTATGAAGCCTATATGTAAGACCTTAAAGAAAG 2
    |||||
```

RESULT 10
US-09-516-448-1391
Sequence 1391, Application US/09516448
GENERAL INFORMATION:
APPLICANT: Robison, Keith E.
TITLE OF INVENTION: Novel Nucleic Acid and Protein Homologs
FILE REFERENCE: 5800-119
CURRENT APPLICATION NUMBER: US/09/516, 448
CURRENT FILING DATE: 2000-02-29
NUMBER OF SEQ ID NOS: 2050
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 1391
LENGTH: 630
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(630)
OTHER INFORMATION: n = A,T,C or G
US-09-516-448-1391

Query Match 74.1%; Score 504.8; DB 19; Length 630;

Best Local Similarity 95.7%; Pred. No. 2e-131;
Matches 603; Conservative 16; Indels 11; Gaps 8;

QY	19	cccaaaacccaactctacatctctccagcgagaaatcggtgtgttttgaagacccttgaagaa	78
Db	2	ccgaaaaaaaccaact-acaatctcccgccgaagaatcggtgtgttttgaagaccctgaagaa	60
QY	79	ctcaaatgtggttttcaactatctgaagagacatctctccagaaatgtggagattgtgaaat	138
Db	61	ctcaatgtggttttcaactatctgaagagacatctctccagaaatgtggagattgtgaaat	120
QY	139	ctagaagagacttggaatttctctggaatctgaatataatgtgagctgccccttgaatga	198
Db	121	ctagaagagagctggaatttctctggaatctgaatataatgtgagctgccccttgaatga	160
QY	199	aatttgaagcagagtataattgtgatatactccacaaacagatttccagtgccccatc	258
Db	181	aatttgaagcagagtataattgtgatatactccacaaacagatttccagtgccccatc	240
QY	259	tgtgtccctgcgagatgcgaatttgcagtggttggatacca-gcagacataacctgaccga	317
Db	241	tgtgtccctgcgagatgcgaatttgcagtggttggatactatgcagacataacctgaccga	300
QY	318	cccttcgcgcagagatatgaacagggcctgacgggagcttcctctctgtgataaaaaca	377
Db	301	cccttcgcgcagagatatgaacagggcctgacgggagcttcctctctgtgataaaaaca	360
QY	378	g-ttgagacttacctcccatctccaatgtgatcaacctggaagagc-tcaactctgttaat-cgt	434
Db	361	gtttgaccttacctcccatctccaatgtgatcaacctggaagagcttcaactctgttaat-cgt	420
QY	435	cagtggggagcca-ttgggtgtgagctcccaca-ctcgcctcttgtgactatccaacact	488
Db	421	camggggagccatttgggtgtgagctcccacaacctgcgcctcttgtgactatccaacac	480
QY	490	ttaaatatttgaagccctatgagacaatcccatgtgataatgcccaatgtgaagatggaat	549
Db	481	ccaaatatttgaagccctatgagacaatcccatgtgataatgcccaatgtgaagatggaat	540
QY	550	gaataatgtgaagatggaac-ggagctgcgccaacatttgaataaagaagttaagaagccta	608
Db	541	gaataatgtgaagatggaagctgcgccaacatttgaataaagaagttaagaagccta	600
QY	609	tattgaagaccttaagaagaagagatctgt 638	
Db	601	tattgaagaccttaagaagaagaagatctgt 630	

```

RESULT 12
US-09-616-111-636
? Sequence 636, Application US/09616111
? GENERAL INFORMATION:
? APPLICANT: Holtzman, Douglas A.
? APPLICANT: Gearing, David P.
? TITLE OF INVENTION: NUCLEIC ACID MOLECULES DERIVED FROM A
? TITLE OF INVENTION: HUMAN PITUITARY GLAND
? FILE REFERENCE: 1600.1142-001
? CURRENT APPLICATION NUMBER: US/09/616,111
? CURRENT FILING DATE: 2000-07-13
? PRIOR APPLICATION NUMBER: 60/143,618
? PRIOR FILING DATE: 1999-07-13
? NUMBER OF SEQ ID NOS: 816
? SOFTWARE: FastSeq for Windows Version 3.0
? SEQ ID NO 636
? LENGTH: 484
? TYPE: DNA
? ORGANISM: Homo sapiens
? FEATURE:
? NAME/KEY: misc_feature
? LOCATION: (1)...(484)

```


US-60-168-611-2255

Query Match

50.7%; Score 345; DB 48; Length 840;

Best Local Similarity 100.0%; Pred. No. 2.6e-86; Mismatches 0; Indels 0; Gaps 0;

Matches 345; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

OY 337 aggtctagagagctgcagagctctctctgtatataaacaagttgactactctccat 396
    |||||||
DB 142 aggtctagagagctgcagagctctctctgtatataaacaagttgactactctccat 201
OY 397 tccatgtcgaacctggaagaagctcactctgttagctcagtcgggacatttgtagag 456
    |||||||
DB 202 tccatgtcgaacctggaagaagctcactctgttagctcagtcgggacatttgtagag 261
OY 457 ctcccaactgcctcttgtagctatccacactttaaatgtgaagccttatgacaat 516
    |||||||
DB 262 ctcccaactgcctcttgtagctatccacactttaaatgtgaagccttatgacaat 321
OY 517 cctatgataatgcccaatgtgaagatggaatgaatgaatgaatgaatgaatgaatgacg 576
    |||||||
DB 322 cctatgataatgcccaatgtgaagatggaatgaatgaatgaatgaatgaatgaatgacg 381
OY 577 caacatttgataagaagatgatgaagcctatactgaagaccttaagaagaagaatct 636
    |||||||
DB 382 caacatttgataagaagatgatgaagcctatactgaagaccttaagaagaagaatct 441
OY 637 gtcccaagctataccaccaagtgctctttagcctcaacttga 681
    |||||||
DB 442 gtcccaagctataccaccaagtgctctttagcctcaacttga 486

```

RESULT 15

US-09-332-782-6811

Sequence 6811, Application US/09332782

GENERAL INFORMATION:

APPLICANT: Hyseq, Inc.

TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED

TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES

FILE REFERENCE: 20411-754CON1

CURRENT APPLICATION NUMBER: US/09/332.782

CURRENT FILING DATE: 1999-06-14

EARLIER APPLICATION NUMBER: US 09/181.430

EARLIER FILING DATE: 1998-10-28

NUMBER OF SEQ ID NOS: 21027

SOFTWARE: FastSeq for Windows Version 3.0

SEQ ID NO 6811

LENGTH: 327

TYPE: DNA

ORGANISM: Homo sapiens

US-09-332-782-6811

Query Match

47.6%; Score 324.4; DB 17; Length 327;

Best Local Similarity 99.7%; Pred. No. 1.2e-80; Mismatches 1; Indels 0; Gaps 0;

Matches 325; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

OY 126 agattgtgaaatctagaagactggtgttctggaatctagaatctagaagctgccc 185
    |||||||
DB 1 agattgtgaaatctagaagactggtgttctggaatctagaatctagaagctgccc 60
OY 186 cttagaattgaattgaagcaagtacattgtatatactcagaacaagtttcc 245
    |||||||
DB 61 cttagaattgaattgaagcaagtacattgtatatactcagaacaagtttcc 120
OY 246 cagtgctccaatctgtctcctcgcgagtcgcaatttcgaatggttgatatacagacgaa 305
    |||||||
DB 121 cagtgctccaatctgtctcctcgcgagtcgcaatttcgaatggttgatatacagacgaa 180
OY 306 taacttaccagactgctgcgcaagaatatacagagctagagagctgcaagacttctct 365
    |||||||
DB 181 taacttaccagactgctgcgcaagaatatacagagctagagagctgcaagacttctct 240

```

```

OY 366 gtataaacaagttgacctactctccctattcctatgctgaacctgaagaagctcaact 425
    |||||||
DB 241 gtataaacaagttgacctactctccctattcctatgctgaacctgaagaagctcaact 300
OY 426 gtagtcgctcagtcggaacatttgg 451
    |||||||
DB 301 gtagtcgctcagtcggaacatttgg 326

```

Search completed: February 26, 2002, 11:47:58
Job time: 7745 sec

```
Db 241 ttctccagttgcccaatctgtctcgtcggaagtgcgaattgtcagtggttgatctcagc 300
|||
Qy 301 agcaataacctgacacgtctgcgcaagataatagacagctagaagagctgcagagcttt 360
|||
Db 301 agcaataacctgacacgtctgcgcaagataatagacagctagaagagctgcagagcttt 360
|||
Qy 361 ctcttgtataaacaagaattgacacttcccttccatctccatctgctgtaaacctgaagaagctc 420
|||
Db 361 ctcttgtataaacaagaattgacacttcccttccatctccatctgctgtaaacctgaagaagctc 420
|||
Qy 421 actctgttaagtcgtcagtgaggagacatttggttgagagctcccaactgcctcttgtaacctca 480
|||
Db 421 actctgttaagtcgtcagtgaggagacatttggttgagagctcccaactgcctcttgtaacctca 480
|||
Qy 481 tccacaccttaaaattgttaagccttaagcaatctcattgtatgaatgcccagaatgtgaa 540
|||
Db 481 tccacaccttaaaattgttaagccttaagcaatctcattgtatgaatgcccagaatgtgaa 540
|||
Qy 541 gatggcaatgaaataatgaaagtgaaagcgatcgccacaatttgataaagaagtatg 600
|||
Db 541 gatggcaatgaaataatgaaagtgaaagcgatcgccacaatttgataaagaagtatg 600
|||
Qy 601 aagagctatattgaaagccttaagaagaagatctgttccagctataccaccaagtgtg 660
|||
Db 601 aagagctatattgaaagccttaagaagaagatctgttccagctataccaccaagtgtg 660
|||
Qy 661 tctttagccttcaacttga 681
|||
Db 661 tctttagccttcaacttga 681
|||
```

```
RESULT 2
US-09-602-833A-1
; Sequence 1, Application US/09602833A
; GENERAL INFORMATION:
; APPLICANT: Turner, Alex
; APPLICANT: Zambrowicz, Brian
; APPLICANT: Nehls, Michael
; APPLICANT: Freidrich, Glenn A.
; APPLICANT: Sands, Arthur T.
; TITLE OF INVENTION: NOVEL HUMAN GENES AND PROTEINS
; TITLE OF INVENTION: ENCODED THEREBY
; FILE REFERENCE: 8535-0036-999
; CURRENT APPLICATION NUMBER: US/09/602,833A
; CURRENT FILING DATE: 2000-06-23
; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 1
; LENGTH: 1116
; TYPE: DNA
; ORGANISM: Homo sapien
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)...(1113)
US-09-602-833A-1
```

```
Query Match 100.0%; Score 681; DB 5; Length 1116;
Best Local Similarity 100.0%; Pred. No. 5, 6e-183;
Matches 681; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 atgagaattctggaatctgcccacaaacccaatctcacatcttcagcagagaatcggtgt 60
|||||
Db 436 atgagaattctggaatctgcccacaaacccaatctcacatcttcagcagagaatcggtgt 495
|||||
Qy 61 ttgagaacctggaagaactcaatggtgttcaactatctggaagagcatccctccagaa 120
|||||
Db 496 ttgagaacctggaagaactcaatggtgttcaactatctggaagagcatccctccagaa 555
|||||
Qy 121 ttggagattctggaataatctgagagagctgattgtcttgtaaatctcgaataatagag 180
|||||
Db 556 ttggagattctggaataatctgagagagctgattgtcttgtaaatctcgaataatagag 615
|||||
```

```
Qy 181 ctgcccttgaattaagtaattgaaagaatttaacatttgtagatctcagcaacaag 240
|||||
Db 616 ctgcccttgaattaagtaattgaaagaatttaacatttgtagatctcagcaacaag 675
|||||
Qy 241 ttctccagtgcccaatctgtgtctcgtcgatgtcgaatttgcaagtggttgatacagc 300
|||||
Db 676 ttctccagtgcccaatctgtgtctcgtcgatgtcgaatttgcaagtggttgatacagc 735
|||||
Qy 301 agcaataacctgacacgtctgcgcaagataatagacagctagaagagctgcagagcttt 360
|||||
Db 736 agcaataacctgacacgtctgcgcaagataatagacagctagaagagctgcagagcttt 795
|||||
Qy 361 ctcttgtataaacaagaattgacacttcccttccatctccatctgctgtaaacctgaagaagctc 420
|||||
Db 796 ctcttgtataaacaagaattgacacttcccttccatctccatctgctgtaaacctgaagaagctc 855
|||||
Qy 421 actctgttaagtcgtcagtgaggagacatttggttgagagctcccaactgcctcttgtaacctca 480
|||
Db 481 actctgttaagtcgtcagtgaggagacatttggttgagagctcccaactgcctcttgtaacctca 915
|||
Qy 481 tccacaccttaaaattgttaagccttaagcaatctcattgtatgaatgcccagaatgtgaa 540
|||
Db 916 tccacaccttaaaattgttaagccttaagcaatctcattgtatgaatgcccagaatgtgaa 975
|||
Qy 541 gatggcaatgaaataatgaaagtgaaagcgatcgccacaatttgataaagaagtatg 600
|||||
Db 976 gatggcaatgaaataatgaaagtgaaagcgatcgccacaatttgataaagaagtatg 1035
|||||
Qy 601 aagagctatattgaaagccttaagaagaagatctgttccagctataccaccaagtgtg 660
|||||
Db 1036 aagagctatattgaaagccttaagaagaagatctgttccagctataccaccaagtgtg 1095
|||||
Qy 661 tctttagccttcaacttga 681
|||
Db 1096 tctttagccttcaacttga 1116
|||
```

```
RESULT 3
US-09-933-524A-29336
; Sequence 29336, Application US/09933524A
; GENERAL INFORMATION:
; APPLICANT: Dimañac, Radoje T.
; APPLICANT: Labat, Ivan
; APPLICANT: Staché-Crain, Birgit
; APPLICANT: Dickson, Mark
; APPLICANT: Jones, Lee W.
; TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
; TITLE OF INVENTION: From Various Libraries
; FILE REFERENCE: 774
; CURRENT APPLICATION NUMBER: US/09/933,524A
; CURRENT FILING DATE: 2001-08-20
; PRIOR APPLICATION NUMBER: 09/528,409
; PRIOR FILING DATE: 2000-05-17
; NUMBER OF SEQ ID NOS: 116231
; SOFTWARE: Hy-patent.pl Version 3.1
; SEQ ID NO 29336
; LENGTH: 480
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-933-524A-29336
```

```
Query Match 25.3%; Score 172; DB 6; Length 480;
Best Local Similarity 100.0%; Pred. No. 1, 2e-38;
Matches 172; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 atgagaattctggaatctgcccacaaacccaatctcacatcttcagcagagaatcggtgt 60
|||||
Db 309 atgagaattctggaatctgcccacaaacccaatctcacatcttcagcagagaatcggtgt 368
|||||
Qy 61 ttgagaacctggaagaactcaatggtgttcaactatctggaagagcatccctccagaa 120
|||||
```

Db 369 ttgaagaacctgaagaactcaatgtggtttcaactatctgaagagcattcctccagaa 428
QY 121 ttggagattgtgaaaaatcagagagactgtgtcttgaaatcagaat 172
Db 429 ttggagattgtgaaaaatcagagagactgtgtcttgaaatcagaat 480

RESULT 4
US-09-933-524A-25490

; Sequence 25490, Application US/09933524A
; GENERAL INFORMATION:
; APPLICANT: Dirmnac, Radoje T.
; APPLICANT: Labat, Ivan
; APPLICANT: Stache-Crain, Birgit
; APPLICANT: Dickson, Mark
; APPLICANT: Jones, Lee W.
; TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
; FILE REFERENCE: 774
; CURRENT APPLICATION NUMBER: US/09/933,524A
; CURRENT FILING DATE: 2001-08-20
; PRIOR APPLICATION NUMBER: 09/528,409
; PRIOR FILING DATE: 2000-05-17
; NUMBER OF SEQ ID NOS: 116231
; SOFTWARE: HY-patent.pl Version 3.1
; SEQ ID NO 25490
; LENGTH: 440
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-933-524A-25490

Query Match 18.8%; Score 128; DB 6; Length 440;
Best Local Similarity 100.0%; Pred. No. 3.7e-26;
Matches 128; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 atgagaattctgattctgcacaaacaaatcacatcttcacagagaatcgtgtg 60
Db 312 atgagaattctgattctgcacaaacaaatcacatcttcacagagaatcgtgtg 371
QY 61 ttgaagaacctgaagaactcaatgtggtttcaactatctgaagagcattcctccagaa 120
Db 372 ttgaagaacctgaagaactcaatgtggtttcaactatctgaagagcattcctccagaa 431
QY 121 ttggagaga 128
Db 432 ttggagaga 439

RESULT 5

US-09-933-524A-24337
; Sequence 24337, Application US/09933524A
; GENERAL INFORMATION:
; APPLICANT: Dirmnac, Radoje T.
; APPLICANT: Labat, Ivan
; APPLICANT: Stache-Crain, Birgit
; APPLICANT: Dickson, Mark
; APPLICANT: Jones, Lee W.
; TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
; FILE REFERENCE: 774
; CURRENT APPLICATION NUMBER: US/09/933,524A
; CURRENT FILING DATE: 2001-08-20
; PRIOR APPLICATION NUMBER: 09/528,409
; PRIOR FILING DATE: 2000-05-17
; NUMBER OF SEQ ID NOS: 116231
; SOFTWARE: HY-patent.pl Version 3.1
; SEQ ID NO 24337
; LENGTH: 456
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature

; LOCATION: (1)...(456)
; OTHER INFORMATION: n = A,T,C or G
US-09-933-524A-24337

Query Match 18.4%; Score 125; DB 6; Length 456;
Best Local Similarity 100.0%; Pred. No. 2.6e-25;
Matches 125; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 atgagaattctgattctgcacaaacaaatcacatcttcacagagaatcgtgtg 60
Db 331 atgagaattctgattctgcacaaacaaatcacatcttcacagagaatcgtgtg 390
QY 61 ttgaagaacctgaagaactcaatgtggtttcaactatctgaagagcattcctccagaa 120
Db 391 ttgaagaacctgaagaactcaatgtggtttcaactatctgaagagcattcctccagaa 450
QY 121 ttggg 125
Db 451 ttggg 455

RESULT 6

US-09-933-524A-13035
; Sequence 13035, Application US/09933524A
; GENERAL INFORMATION:
; APPLICANT: Dirmnac, Radoje T.
; APPLICANT: Labat, Ivan
; APPLICANT: Stache-Crain, Birgit
; APPLICANT: Dickson, Mark
; APPLICANT: Jones, Lee W.
; TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
; FILE REFERENCE: 774
; CURRENT APPLICATION NUMBER: US/09/933,524A
; CURRENT FILING DATE: 2001-08-20
; PRIOR APPLICATION NUMBER: 09/528,409
; PRIOR FILING DATE: 2000-05-17
; NUMBER OF SEQ ID NOS: 116231
; SOFTWARE: HY-patent.pl Version 3.1
; SEQ ID NO 13035
; LENGTH: 429
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-933-524A-13035

Query Match 12.8%; Score 87; DB 6; Length 429;
Best Local Similarity 100.0%; Pred. No. 1.6e-14;
Matches 87; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 atgagaattctgattctgcacaaacaaatcacatcttcacagagaatcgtgtg 60
Db 343 atgagaattctgattctgcacaaacaaatcacatcttcacagagaatcgtgtg 402
QY 61 ttgaagaacctgaagaactcaatgtg 87
Db 403 ttgaagaacctgaagaactcaatgtg 429

RESULT 7

US-09-933-524A-15579
; Sequence 15579, Application US/09933524A
; GENERAL INFORMATION:
; APPLICANT: Dirmnac, Radoje T.
; APPLICANT: Labat, Ivan
; APPLICANT: Stache-Crain, Birgit
; APPLICANT: Dickson, Mark
; APPLICANT: Jones, Lee W.
; TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
; FILE REFERENCE: 774
; CURRENT APPLICATION NUMBER: US/09/933,524A

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?
?
? ORGANISM: Homo sapiens
?
? OS-09-933-524A-15579
?
?
?
?
? CURRENT FILING DATE: 2001-08-20
?
? PRIOR APPLICATION NUMBER: 09/528,409
?
?
? PRIOR FILING DATE: 2000-05-17
?
? NUMBER OF SEQ ID NOS: 116231
?
? SOFTWARE: HyPatent.pl Version 3.1
?
? SEQ ID NO 15579
?
? LENGTH: 428
?
? TYPE: DNA
?

```

Query Match	11.48;	Score 77.6;	DB 6;	Length 428;
Best Local Similarity	50.58;	Pred. No. 7.3e-12;		
Matches 188; Conservative	0;	Mismatches 184;	Indels 0;	Gaps 0;

QY	91	ttcaacatctcgaagagcatccctcccgaaattggagagattgtaaaatcagaggagatg	150
Db	21	tcgacaacaatcaagactgtccccaagaaactaagtattgtgtccagcttgggaaacta	80
QY	151	gattgtctctgnaaactcagaatlaatgaatgagctgccttgaatgaatlaattggaacaa	210
Db	81	gaactggtctgttaacagagatatatgtaacttccacaagagctcagcaatctgtcaaa	140
QY	211	gttaacatttgaatatactctcagcaaaagtatttccagtggtcccaatctgtctctgcgg	270
Db	141	cttactacaactgtactctgagatgaatgaacgatttactacaactccctctgtgtgtgaac	200
QY	271	atgfcgcaatttgcagtggttggaatcagacgaacataactgcagccgcgcgcgacagat	330
Db	201	atgcccgtcccttgggtgtgcggaatvgggaagcaacaactgtgaacaacttccgtgaact	260
QY	331	atagacaggtcgaaggagcgtgcagagcttctctctgtatataaaacaagtgcactacct	390
Db	261	atagaaaagaatgcacaactctcaatcagcttatgtgtgaagaagaatgaataaactgttg	320
QY	391	cccatctccatgctgaaacctgaaagagctcaactctgtttagtcgacgtggggacattg	450
Db	321	ccctaaacaactcagcatalatgaaaaaactgtgtactcttgttctcagcaacaataaactg	380
QY	451	gtggagctccca	462
Db	381	caagataatccca	392

```
RESULT      8
US-09-898-888A-13202
Sequence 13202, Application US/098698888A
GENERAL INFORMATION:
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL CONTIGS OBTAINED FROM VARIOUS CDNAs
TITLE OF INVENTION: LIBRARIES
FILE REFERENCE: 20411-748CON1
CURRENT APPLICATION NUMBER: US/09/898,888A
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/340,623
PRIOR FILING DATE: 1999-06-28
PRIOR APPLICATION NUMBER: US 09/205,070
PRIOR FILING DATE: 1998-12-03
NUMBER OF SEQ ID NOS: 45207
SOFTWARE: FASTSeq for Windows Version 3.0
SEQ ID NO 13202
LENGTH: 1936
TYPE: DNA
ORGANISM: Homo sapiens
US-09-898-888A-13202
```

Query Match	8.4%;	Score 57.4;	DB 6;	Length 1936;
Best Local Similarity	47.0%;	Pred. No. 7.5e-06;		
Matches 178; Conservative	0;	Mismatches 201;	Indels 0;	Gaps 0;

182 tgccttgaatgaagtaattgaagcaagttacattgtgatatctcagcaacaagt 241

Db	1167	taccagcggagcattttagtttaccagaactccagatgcttagatgttgagctaacaca	1226
Qy	242	tttcccgatgtcccaatcctgtgtccctgcgagatgcgaatttcagtcagtcgttggatatcagca	301
Db	1227	tttcaatgatctccaatagaatgagattgtcttcagaacctgcgaatttgcatactatcg	1286
Qy	302	gcaaatcccgaccgcacccgcagatatagacagcgttagaagcagtcgcagccttc	361
Db	1287	ggacaacaagcggagcattcttcgccaacaactgttttaagcatgaagtcgttgagcttga	1346
Qy	362	tcttgtataaacaacagctgcacctacccttcctattccatgcgtcaacctgaagaagctca	421
Db	1347	atctcgagacacactgcatactaccctccaccagaagaagtgtgtaagcttccaccgctca	1406
Qy	422	ctctggttagcgtccagctggagaccatttggctggagctcccaactgccttttggctcat	481
Db	1407	ctcagctcgagcgtcgaagcgggaactgcttgcagcgcctcgacacccagcttggccagtc	1466
Qy	482	ccacaccccttaaaatttgcgaagccttatgcacatccctatgcataagcccaatgtgaag	541
Db	1467	ggatgcctcagaagaacgcggcctgtgttgcgaagatcaccttltatatacctgcacatcg	1526
Qy	542	atgcgaatgaataatgcga	560
Db	1527	aagtcacaagcagcatcgaa	1515

```

RESULT      9
US-09-902-775A-184
: Sequence 184, Application US/09902775A
: GENERAL INFORMATION:
: APPLICANT: Genentech, Inc.
: APPLICANT: Ashkenazi, Avi
: APPLICANT: Botstein, David
: APPLICANT: Desnoyers, Luc
: APPLICANT: Eaton, Dan L.
: APPLICANT: Ferrara, Napoleone
: APPLICANT: Filvaroff, Ellen
: APPLICANT: Fong, Sherman
: APPLICANT: Gao, Wei-Qiang
: APPLICANT: Gerber, Hanspeter
: APPLICANT: Gerltsen, Mary E.
: APPLICANT: Goddard, A.
: APPLICANT: Godowski, Paul J.
: APPLICANT: Grimaldi, Christopher J.
: APPLICANT: Gurney, Austin L.
: APPLICANT: Hillan, Kenneth, J.
: APPLICANT: Kijavlin, Ivar J.
: APPLICANT: Mather, Jennie P.
: APPLICANT: Pan, James
: APPLICANT: Paoni, Nicholas F.
: APPLICANT: Roy, Margaret Ann
: APPLICANT: Stewart, Timothy A.
: APPLICANT: Tumas, Daniel
: APPLICANT: Williams, P. Mickey
: APPLICANT: Wood, William, I.
: TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
: TITLE OF INVENTION: Acids Encoding the Same
: FILE REFERENCE: 10466-14
: CURRENT APPLICATION NUMBER: US/09/902,775A
: CURRENT FILING DATE: 2001-07-10
: PRIOR APPLICATION NUMBER: PCT/US00/04414
: PRIOR FILING DATE: 2000-02-22
: PRIOR APPLICATION NUMBER: US 60/143,048
: PRIOR FILING DATE: 1999-07-07
: PRIOR APPLICATION NUMBER: US 60/145,698
: PRIOR FILING DATE: 1999-07-26
: PRIOR APPLICATION NUMBER: US 60/146,222
: PRIOR FILING DATE: 1999-07-28
: PRIOR APPLICATION NUMBER: PCT/US99/20594
: PRIOR FILING DATE: 1999-09-08
: PRIOR APPLICATION NUMBER: PCT/US99/20944

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; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 184
; LENGTH: 1947
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-902-775A-184
```

```

Query Match      8.4%; Score 57.4; DB 5; Length 1947;
Best Local Similarity 47.0%; Pred. No. 7.5e-06;
Matches 178; Conservative 0; Mismatches 201; Indels 0; Gaps 0;
```

```

QY 182 tgccttgaatgaatgaatgaagcaagttacattgttagatatcagaacaaga 241
    || || || || || || || || || || || || || || || || || || ||
DB 1162 taccagtgagcagattagttacagaactcagatgctagtgagctaacaaaca 1221
    || || || || || || || || || || || || || || || || || || ||
QY 242 ttccagtgcccaatctgtctcctgcgagtgcgaattgcgaagtgttgagatacga 301
    || || || || || || || || || || || || || || || || || || ||
DB 1222 ttccatgatcccaatagaatagatgtcttcgaactgcagcatcttgcatatcactg 1281
    || || || || || || || || || || || || || || || || || || ||
QY 302 gcaataaccctgacccgacccaagatagacagcgtagagcgagctgcagaagcttc 361
    || || || || || || || || || || || || || || || || || || ||
DB 1282 ggaacaagaatgacatcttcgccaacaactgtttaagcataaagttgaggaacttga 1341
    || || || || || || || || || || || || || || || || || || ||
QY 362 tcttgataaacaagaatgtgacctacatccatcattcattgctgaactgaaagactca 421
    || || || || || || || || || || || || || || || || || || ||
DB 1342 atctggagacagactcattcaccctcaccagagaagttgtgtcagctctccagctca 1401
    || || || || || || || || || || || || || || || || || || ||
QY 422 ctctgttagtcgtcagtgaggacatttggtagctcccaactgccttgtgactcat 481
    || || || || || || || || || || || || || || || || || || ||
DB 1402 cttagctgagctgaaggggaactgtcttgaccgctgcagccagctggggcagtgctc 1461
    || || || || || || || || || || || || || || || || || || ||
QY 482 ccacaccccttaaatgtgaagccttatgacacatcctcttaagtaagcccaatggaag 541
    || || || || || || || || || || || || || || || || || || ||
DB 1462 ggaatgctcaagaagaagcgggcttgtgtggaagatacactttttagaccctgccaactg 1521
    || || || || || || || || || || || || || || || || || || ||
QY 542 atggaatgaataatgga 560
    || || || || || || || || || || || || || || || || || || ||
DB 1522 aagtcagaagagcattgaa 1540
    || || || || || || || || || || || || || || || || || || ||
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RESULT 10
US-09-904-956-184
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; Sequence 184, Application US/09904956
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
```

```

; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerltzen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavlin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumasi, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/904,956
; CURRENT FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 184
; LENGTH: 1947
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-904-956-184
```

```

Query Match      8.4%; Score 57.4; DB 5; Length 1947;
Best Local Similarity 47.0%; Pred. No. 7.5e-06;
Matches 178; Conservative 0; Mismatches 201; Indels 0; Gaps 0;
```

```

QY 182 tgccttgaatgaatgaatgaagcaagttacattgttagatatcagaacaaga 241
    || || || || || || || || || || || || || || || || || || ||
```


	Query Match	Similarity	8.4%	Score 57.4	DB 5	Length 1947
	Best Local	Similarity	47.0%	Pred. No. 7.5e-06		
	Matches	178	Conservative	0	Mismatches 201	Indels 0
						Gaps 0
OY	182	tgcccttgaattaaagtaattggaagcaagttacattgtgaatatctgacaaacagt	241			
Ob	1162	taccagttgacgattcttggctttacaggaactaagctgtgcttaattgtgacacaca	1221			

```

1 RESULT 13
2 US-09-909-204-184
3 Sequence 184, Application US/09390204
4 GENERAL INFORMATION:
5 APPLICANT: Genentech, Inc.
6 APPLICANT: Ashkenazi, Avi
7 APPLICANT: Botstein, David
8 APPLICANT: Desnoyers, Luc
9 APPLICANT: Eaton, Dan L.
10 APPLICANT: Ferrara, Napoleone
11 APPLICANT: Filvaroff, Ellen
12 APPLICANT: Fong, Sherman
13 APPLICANT: Gao, Wei-Qiang
14 APPLICANT: Gerber, Hanspeter
15 APPLICANT: Gerlitsen, Mary E.
16 APPLICANT: Goddard, A.
17 APPLICANT: Godowski, Paul J.
18 APPLICANT: Grimaldi, Christopher J.
19 APPLICANT: Gurney, Austin L.
20 APPLICANT: Hillan, Kenneth, J.
21 APPLICANT: Kijavlin, Ivar J.
22 APPLICANT: Mather, Jennie P.
23 APPLICANT: Pan, James
24 APPLICANT: Peoni, Nicholas F.
25 APPLICANT: Roy, Margaret Ann
26 APPLICANT: Stewart, Timothy A.
27 APPLICANT: Tumas, Daniel
28 APPLICANT: Williams, P. Mickey
29 APPLICANT: Wood, William, I.
30 TITLE OF INVENTION: Acids Encoding the Same
31 TITLE OF INVENTION: Acids Encoding the Same
32 FILE REFERENCE: 10466-14
33 CURRENT APPLICATION NUMBER: US/09/909,204
34 CURRENT FILING DATE: 2001-07-18
35 PRIOR APPLICATION NUMBER: PCT/US00/04414
36 PRIOR FILING DATE: 2000-02-22
37 PRIOR APPLICATION NUMBER: US 60/143,048
38 PRIOR FILING DATE: 1999-07-07
39 PRIOR APPLICATION NUMBER: US 60/145,698
40 PRIOR FILING DATE: 1999-07-26
41 PRIOR APPLICATION NUMBER: US 60/146,222
42 PRIOR FILING DATE: 1999-07-28
43 PRIOR APPLICATION NUMBER: PCT/US99/20594
44 PRIOR FILING DATE: 1999-09-08
45 PRIOR APPLICATION NUMBER: PCT/US99/20944
46 PRIOR FILING DATE: 1999-09-13
47 PRIOR APPLICATION NUMBER: PCT/US99/21090

```



```
QY 242 ttccagtgcccaatctgtctcgtcgatgctgaattgcagtggttgatatacga 301
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1222 ttccatgattccaataagaattagattgcttcaagaaccgcagcattgacatacctg 1281
QY 302 gcaataactgaccagcctgcgcgaagatatagaagcgttagaggtgcagaagcttc 361
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1282 ggaacaagaaggagcattctcccaaaacaaattgtttaaatgcataaagttagagcttga 1341
QY 362 tctgtatataaacaagttgacctacacctccattccatccatgcagcctgaactgaagaagtc 421
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1342 atctggagcaggaactgcatacctcaccctccagagaagtgtgtcagctctccagctca 1401
QY 422 ctctgttagctgcagtgaggagcattgtgtgagctcccaactgcacctgtgactcat 481
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1402 ctcaagctgagcgtcgaaggaactgctgtgacccgcctgcacgcccagcttgagcaggtgc 1461
QY 482 ccacaccttaaatgtgtaagccttatggaacaactctattgataatgtgcaatgtgaag 541
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1462 ggaatgtcagaagaacgcggctgtgtgtggaagatacaccttctgataccctgcagctcg 1521
QY 542 atggcaatgaataatga 560
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1522 aagtcagaagagcattgaa 1540
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
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```
RESULT 15
US-09-902-634A-184
; Sequence 184, Application US/09902634A
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvarolf, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerltzen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kijavlin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/902,634A
; CURRENT FILING DATE: 2001-07-10
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
```

```
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 184
; LENGTH: 1947
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-902-634A-184
```

```
Query Match 8.48; Score 57.4; DB 5; Length 1947;
Best Local Similarity 47.0%; Pred. No. 7.5e-06;
Matches 178; Conservative 0; Mismatches 201; Indels 0; Gaps 0;
```

```
QY 182 tgccttgtaataaagttgaagcaagttacatttgatagatctcgcaaaagtc 241
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1162 taccagtgagcaatttagttacagaacccaagtccttagatgagctacacaca 1221
QY 242 ttccagtgcccaatctgtctcgtcgatgctgaattgcaagtggttgatatacga 301
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1222 ttcaatgattccaataagaattagattgcttcaagaaccgcagcattgacatacctg 1281
QY 302 gcaataactgaccagcctgcgcgaagatatagaagcgttagaggtgcagaagcttc 361
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1282 ggaacaagaaggagcattctcccaaaacaaattgtttaaatgcataaagttagagcttga 1341
QY 362 tctgtatataaacaagttgacctacacctccattccatccatgcagcctgaactgaagaagtc 421
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1342 atctggagcaggaactgcatacctcaccctccagagaagtgtgtcagctctccagctca 1401
QY 422 ctctgttagctgcagtgaggagcattgtgtgagctcccaactgcacctgtgactcat 481
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1402 ctcaagctgagcgtcgaaggaactgctgtgacccgcctgcacgcccagcttgagcaggtgc 1461
QY 482 ccacaccttaaatgtgtaagccttatggaacaactctattgataatgtgcaatgtgaag 541
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1462 ggaatgtcagaagaacgcggctgtgtgtggaagatacaccttctgataccctgcagctcg 1521
QY 542 atggcaatgaataatga 560
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1522 aagtcagaagagcattgaa 1540
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
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Search completed: February 26, 2002, 11:57:08
Job time: 7170 sec
